

What Is Claimed Is:

1. A protease inhibitor comprising the sequence:

X<sup>1</sup>-Val-Cys-Ser-Glu-Gln-Ala-Glu-X<sup>2</sup>-Gly-X<sup>3</sup>-  
Cys-Arg-Ala-X<sup>4</sup>-X<sup>5</sup>-X<sup>6</sup>-X<sup>7</sup>-Trp-Tyr-Phe-Asp-  
Val-Thr-Glu-Gly-Lys-Cys-Ala-Pro-Phe-X<sup>8</sup>-  
Tyr-Gly-Gly-Cys-X<sup>9</sup>-X<sup>10</sup>-X<sup>11</sup>-X<sup>12</sup>-Asn-Asn-Phe-  
Asp-Thr-Glu-Glu-Tyr-Cys-Met-Ala-Val-Cys-  
Gly-Ser-Ala-Ile,

wherein:

X<sup>1</sup> is selected from Glu-Val-Val-Arg-Glu-, Asp, or Glu;

X<sup>2</sup> is selected from Thr, Val, Ile and Ser;

X<sup>3</sup> is selected from Pro and Ala;

X<sup>4</sup> is selected from Arg, Ala, Leu, Gly, or Met;

X<sup>5</sup> is selected from Ile, His, Leu, Lys, Ala, or Phe;

X<sup>6</sup> is selected from Ser, Ile, Pro, Phe, Tyr, Trp, Asn, Leu, His, Lys, or Glu;

X<sup>7</sup> is selected from Arg, His, or Ala;

X<sup>8</sup> is selected from Phe, Val, Leu, or Gly;

X<sup>9</sup> is selected from Gly, Ala, Lys, Pro, Arg, Leu, Met, or Tyr;

X<sup>10</sup> is selected from Ala, Arg, or Gly;

X<sup>11</sup> is selected from Lys, Ala, or Asn;

X<sup>12</sup> is selected from Ser, Ala, or Arg;

provided that:

when X<sup>4</sup> is Arg, X<sup>6</sup> is Ile;

when X<sup>9</sup> is Arg, X<sup>4</sup> is Ala or Leu; when X<sup>9</sup> is Tyr, X<sup>4</sup> is Ala or X<sup>5</sup> is His; and

either X<sup>5</sup> is not Ile; or X<sup>6</sup> is not Ser; or X<sup>9</sup> is not Leu, Phe, Met, Tyr, or Asn; or X<sup>10</sup> is not Gly; or X<sup>11</sup> is not Asn; or X<sup>12</sup> is not Arg.

2. A protease inhibitor comprising the sequence:

X<sup>1</sup>-Val-Cys-Ser-Glu-Gln-Ala-Glu-Thr-Gly-  
Pro-Cys-Arg-Ala-X<sup>2</sup>-X<sup>3</sup>-X<sup>4</sup>-Arg-Trp-Tyr-Phe-  
Asp-Val-Thr-Glu-Gly-Lys-Cys-Ala-Pro-Phe-  
Phe-Tyr-Gly-Gly-Cys-X<sup>5</sup>-Gly-Asn-Arg-Asn-

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Asn-Phe-Asp-Thr-Glu-Glu-Tyr-Cys-Met-Ala-  
Val-Cys-Gly-Ser-Ala-Ile,

wherein:

X<sup>1</sup> is selected from Glu-Val-Val-Arg-Glu-, Asp, or Glu;

X<sup>2</sup> is selected from Ala, Leu, Gly, or Met;

X<sup>3</sup> is selected from Ile, His, Leu, Lys, Ala, or Phe;

X<sup>4</sup> is selected from Ser, Ile, Pro, Phe, Tyr, Trp, Asn, Leu, His, Lys, or Glu;

X<sup>5</sup> is selected from Gly, Ala, Lys, Pro, Arg, Leu, Met, or Tyr;

provided that:

when X<sup>5</sup> is Arg, X<sup>2</sup> is Ala or Leu; when X<sup>5</sup> is Tyr, X<sup>2</sup> is Ala or X<sup>3</sup> is His; and

either X<sup>3</sup> is not Ile; or X<sup>4</sup> is not Ser; or X<sup>5</sup> is not Leu, Phe, Met, Tyr, or Asn.

3. A protease inhibitor comprising the sequence:

Glu-Val-Val-Arg-Glu-Val-Cys-Ser-Glu-Gln-  
Ala-Glu-Thr-Gly-Pro-Cys-Arg-Ala-X<sup>1</sup>-X<sup>2</sup>-X<sup>3</sup>-  
Arg-Trp-Tyr-Phe-Asp-Val-Thr-Glu-Gly-Lys-  
Cys-Ala-Pro-Phe-Phe-Tyr-Gly-Gly-Cys-X<sup>4</sup>-  
Gly-Asn-Arg-Asn-Asn-Phe-Asp-Thr-Glu-Glu-  
Tyr-Cys-Met-Ala-Val-Cys-Gly-Ser-Ala-Ile,

wherein:

X<sup>1</sup> is selected from Ala, Leu, Gly, or Met;

X<sup>2</sup> is selected from Ile, His, Leu, Lys, Ala, or Phe;

X<sup>3</sup> is selected from Ser, Ile, Pro, Phe, Tyr, Trp, Asn, Leu, His, Lys, or Glu;

X<sup>4</sup> is selected from Gly, Arg, Leu, Met, or Tyr;

provided that:

when X<sup>1</sup> is Ala, X<sup>2</sup> is Ile, His, or Leu;

when X<sup>1</sup> is Leu, X<sup>2</sup> is Ile or His;

when X<sup>1</sup> is Leu and X<sup>2</sup> is Ile, X<sup>3</sup> is not Ser;

when X<sup>1</sup> is Gly, X<sup>2</sup> is Ile;

when X<sup>4</sup> is Arg, X<sup>1</sup> is Ala or Leu;

when X<sup>4</sup> is Tyr, X<sup>1</sup> is Ala or X<sup>2</sup> is His; and

either X<sup>1</sup> is not Met, or X<sup>2</sup> is not Ile, or X<sup>3</sup> is not Ser, or X<sup>4</sup> is not Gly.

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4. A protease inhibitor according to claim 1, wherein at least two amino acid residues selected from the group consisting of X<sup>4</sup>, X<sup>5</sup>, X<sup>6</sup>, and X<sup>7</sup> differ from the residues found in the naturally occurring sequence of KPI.

5. A protease inhibitor according to claim 1, wherein X<sup>1</sup> is Asp or Glu, X<sup>2</sup> is Thr, X<sup>3</sup> is Pro, and X<sup>12</sup> is Ser.

6. A protease inhibitor according to claim 5, wherein X<sup>1</sup> is Glu, X<sup>2</sup> is Thr, X<sup>3</sup> is Pro, X<sup>4</sup> is Met, X<sup>5</sup> is Ile, X<sup>6</sup> is Ser, X<sup>7</sup> is Arg, X<sup>8</sup> is Phe, X<sup>9</sup> is Gly, X<sup>10</sup> is Gly, and X<sup>11</sup> is Asn.

7. A protease inhibitor according to claim 5, wherein X<sup>1</sup> is Asp, X<sup>2</sup> is Thr, X<sup>3</sup> is Pro, X<sup>4</sup> is Arg, X<sup>5</sup> is Ile, X<sup>6</sup> is Ile, X<sup>7</sup> is Arg, X<sup>8</sup> is Val, X<sup>9</sup> is Arg, X<sup>10</sup> is Ala, and X<sup>11</sup> is Lys.

8. A protease inhibitor according to claim 1, wherein X<sup>1</sup> is Glu-Val-Val-Arg-Glu-, X<sup>2</sup> is Thr, X<sup>3</sup> is Pro, X<sup>4</sup> is Met, X<sup>5</sup> is Ile, X<sup>6</sup> is Ser, X<sup>7</sup> is Arg, X<sup>8</sup> is Phe, X<sup>9</sup> is Gly, X<sup>10</sup> is Gly, X<sup>11</sup> is Asn, and X<sup>12</sup> is Ala.

9. A protease inhibitor according to claim 1, wherein X<sup>1</sup> is Glu-Val-Val-Arg-Glu-, X<sup>2</sup> is Thr, X<sup>3</sup> is Pro, X<sup>4</sup> is Met, X<sup>5</sup> is Ile, X<sup>6</sup> is Ser, X<sup>7</sup> is Arg, X<sup>8</sup> is Phe, X<sup>9</sup> is Gly, X<sup>10</sup> is Gly, X<sup>11</sup> is Ala, and X<sup>12</sup> is Arg.

10. A protease inhibitor according to claim 1, wherein X<sup>1</sup> is Glu, X<sup>2</sup> is Thr, X<sup>3</sup> is Pro, X<sup>4</sup> is Met, X<sup>5</sup> is Ile, X<sup>6</sup> is Ser, X<sup>7</sup> is Arg, X<sup>8</sup> is Phe, X<sup>9</sup> is Gly, X<sup>10</sup> is Ala, X<sup>11</sup> is Asn, and X<sup>12</sup> is Arg.

11. A protease inhibitor according to claim 1, wherein X<sup>1</sup> is Glu-Val-Val-Arg-Glu-, X<sup>2</sup> is Thr, X<sup>3</sup> is Pro, X<sup>4</sup> is Met, X<sup>5</sup> is Ile, X<sup>6</sup> is Ser, X<sup>7</sup> is Arg, X<sup>8</sup> is Phe, X<sup>9</sup> is Gly, X<sup>10</sup> is Arg, X<sup>11</sup> is Asn, and X<sup>12</sup> is Arg.

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12. A protease inhibitor according to claim 1, wherein X<sup>1</sup> is Glu-Val-Val-Arg-Glu-, X<sup>2</sup> is Thr, X<sup>3</sup> is Pro, X<sup>4</sup> is Met, X<sup>5</sup> is Ile, X<sup>6</sup> is Ser, X<sup>7</sup> is Arg, X<sup>8</sup> is Val, Leu, or Gly, X<sup>9</sup> is Gly, X<sup>10</sup> is Gly, X<sup>11</sup> is Asn, and X<sup>12</sup> is Arg.

13. A protease inhibitor according to claim 1, wherein X<sup>1</sup> is Glu-Val-Val-Arg-Glu-, X<sup>2</sup> is Thr, X<sup>3</sup> is Pro, X<sup>4</sup> is Met, X<sup>5</sup> is Ile, X<sup>6</sup> is Ser, X<sup>7</sup> is Ala, X<sup>8</sup> is Phe, X<sup>9</sup> is Gly, X<sup>10</sup> is Gly, X<sup>11</sup> is Asn, and X<sup>12</sup> is Arg.

14. A protease inhibitor according to claim 1, wherein X<sup>1</sup> is Glu-Val-Val-Arg-Glu-, X<sup>2</sup> is Thr, Val, or Ser, X<sup>3</sup> is Pro, X<sup>4</sup> is Ala or Leu, X<sup>5</sup> is Ile, X<sup>6</sup> is Tyr, X<sup>7</sup> His, X<sup>8</sup> is Phe, X<sup>9</sup> is Gly, X<sup>10</sup> is Gly, X<sup>11</sup> is Ala, and X<sup>12</sup> is Arg.

15. A protease inhibitor according to claim 14, wherein X<sup>2</sup> is Thr, and X<sup>4</sup> is Ala.

16. A protease inhibitor according to claim 14, wherein X<sup>2</sup> is Thr, and X<sup>4</sup> is Leu.

17. A protease inhibitor according to claim 14, wherein X<sup>2</sup> is Val, and X<sup>4</sup> is Ala.

18. A protease inhibitor according to claim 14, wherein X<sup>2</sup> is Ser, and X<sup>4</sup> is Ala.

19. A protease inhibitor according to claim 14, wherein X<sup>2</sup> is Val, and X<sup>4</sup> is Leu.

20. A protease inhibitor according to claim 14, wherein X<sup>2</sup> is Ser, and X<sup>4</sup> is Leu.

21. A protease inhibitor according to claim 1, wherein X<sup>1</sup> is Glu-Val-Val-Arg-Glu-, X<sup>2</sup> is Thr, X<sup>3</sup> is Pro, X<sup>4</sup> is Leu, X<sup>5</sup> is Phe, X<sup>6</sup> is Lys, X<sup>7</sup> is Arg, X<sup>8</sup> is Phe, X<sup>9</sup> is Gly, X<sup>10</sup> is Gly, X<sup>11</sup> is Ala, and X<sup>12</sup> is Arg.

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22. A protease inhibitor according to claim 1, wherein X<sup>1</sup> is Glu-Val-Val-Arg-Glu-, X<sup>2</sup> is Thr, X<sup>3</sup> is Pro, X<sup>4</sup> is Leu, X<sup>5</sup> is Phe, X<sup>6</sup> is Lys, X<sup>7</sup> is Arg, X<sup>8</sup> is Phe, X<sup>9</sup> is Tyr, X<sup>10</sup> is Gly, X<sup>11</sup> is Ala, and X<sup>12</sup> is Arg.

23. A protease inhibitor according to claim 1, wherein X<sup>1</sup> is Glu-Val-Val-Arg-Glu-, X<sup>2</sup> is Thr, X<sup>3</sup> is Pro, X<sup>4</sup> is Leu, X<sup>5</sup> is Phe, X<sup>6</sup> is Lys, X<sup>7</sup> is Arg, X<sup>8</sup> is Phe, X<sup>9</sup> is Leu, X<sup>10</sup> is Gly, X<sup>11</sup> is Ala, and X<sup>12</sup> is Arg.

24. A protease inhibitor according to claim 2, wherein X<sup>1</sup> is Glu, X<sup>2</sup> is Met, X<sup>3</sup> is Ile, X<sup>4</sup> is Ile, and X<sup>5</sup> is Gly.

25. A protease inhibitor according to claim 3, wherein X<sup>1</sup> is Met, X<sup>3</sup> is Ser, and X<sup>4</sup> is Gly.

26. A protease inhibitor according to claim 25, wherein X<sup>2</sup> is selected from His, Ala, Phe, Lys, and Leu.

27. A protease inhibitor according to claim 26, wherein X<sup>2</sup> is His.

28. A protease inhibitor according to claim 27, wherein X<sup>2</sup> is Ala.

29. A protease inhibitor according to claim 27, wherein X<sup>2</sup> is Phe.

30. A protease inhibitor according to claim 27, wherein X<sup>2</sup> is Lys.

31. A protease inhibitor according to claim 27, wherein X<sup>2</sup> is Leu.

32. A protease inhibitor according to claim 3, wherein X<sup>1</sup> is Met, X<sup>2</sup> is Ile, and X<sup>4</sup> is Gly.

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33. A protease inhibitor according to claim 32, wherein X<sup>3</sup> is Ile.

34. A protease inhibitor according to claim 32, wherein X<sup>3</sup> is Pro.

35. A protease inhibitor according to claim 32, wherein X<sup>3</sup> is Phe.

36. A protease inhibitor according to claim 32, wherein X<sup>3</sup> is Tyr.

37. A protease inhibitor according to claim 32, wherein X<sup>3</sup> is Trp.

38. A protease inhibitor according to claim 32, wherein X<sup>3</sup> is Asn.

39. A protease inhibitor according to claim 32, wherein X<sup>3</sup> is Leu.

40. A protease inhibitor according to claim 32, wherein X<sup>3</sup> is Lys.

41. A protease inhibitor according to claim 32, wherein X<sup>3</sup> is His.

42. A protease inhibitor according to claim 32, wherein X<sup>3</sup> is Glu.

43. A protease inhibitor according to claim 3, wherein X<sup>1</sup> is Ala.

44. A protease inhibitor according to claim 43, wherein X<sup>2</sup> is Ile.

45. A protease inhibitor according to claim 44, wherein X<sup>3</sup> is Phe, and X<sup>4</sup> is Gly.

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46. A protease inhibitor according to claim 44, wherein X<sup>3</sup> is Tyr, and X<sup>4</sup> is Gly.

47. A protease inhibitor according to claim 44, wherein X<sup>3</sup> is Trp, and X<sup>4</sup> is Gly.

48. A protease inhibitor according to claim 44, wherein X<sup>3</sup> is Ser or Phe, and X<sup>4</sup> is Arg or Tyr.

49. A protease inhibitor according to claim 43, wherein X<sup>2</sup> is His or Leu, X<sup>3</sup> is Phe, and X<sup>4</sup> is Gly.

50. A protease inhibitor according to claim 3, wherein X<sup>1</sup> is Leu.

51. A protease inhibitor according to claim 50, wherein X<sup>2</sup> is His, X<sup>3</sup> is Asn or Phe, and X<sup>4</sup> is Gly.

52. A protease inhibitor according to claim 50, wherein X<sup>2</sup> is Ile, X<sup>3</sup> is Pro, and X<sup>4</sup> is Gly.

53. A protease inhibitor according to claim 3, wherein X<sup>1</sup> is Gly, X<sup>2</sup> is Ile, X<sup>3</sup> is Tyr, and X<sup>4</sup> is Gly.

54. A protease inhibitor according to claim 3, wherein X<sup>1</sup> is Met, X<sup>2</sup> is His, X<sup>3</sup> is Ser, and X<sup>4</sup> is Tyr.

55. An isolated DNA molecule comprising a DNA sequence encoding a protease inhibitor according to claim 1.

56. An isolated DNA molecule according to claim 55, operably linked to a regulatory sequence that controls expression of the coding sequence in a host cell.

57. An isolated DNA molecule according to claim 56, further comprising a DNA sequence encoding a secretory signal peptide. .

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58. An isolated DNA molecule according to claim 57, wherein said secretory signal peptide comprises the signal sequence of yeast alpha-mating factor.

59. A host cell transformed with a DNA molecule according to claim 55.

60. A host cell according to claim 59, wherein said host cell is *E. coli* or a yeast cell.

61. A host cell according to claim 60, wherein said host cell is *Saccharomyces cerevisiae*.

62. A method for producing a protease inhibitor, comprising the steps of culturing a host cell according to claim 59 and isolating and purifying said protease inhibitor.

63. A pharmaceutical composition, comprising a protease inhibitor according to claim 1, together with a pharmaceutically acceptable sterile vehicle.

64. A method of treatment of a clinical condition associated with increased activity of one or more serine proteases, comprising administering to a patient suffering from said clinical condition an effective amount of a pharmaceutical composition according to claim 63.

65. The method of treatment of claim 64, wherein said clinical condition is blood loss during surgery.

66. A method for inhibiting the activity of serine proteases of interest in a mammal comprising administering a therapeutically effective dose of a pharmaceutical composition according to claim 63.

67. The method of claim 66, wherein said serine proteases are selected from the group consisting of:

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kallikrein; chymotrypsins A and B; trypsin; elastase; subtilisin; coagulants and procoagulants, particularly those in active form, including coagulation factors such as factors VIIa, IXa, Xa, XIa, and XIIa; plasmin; thrombin; proteinase-3; enterokinase; acrosin; cathepsin; urokinase; and tissue plasminogen activator.

68. A protease inhibitor comprising the sequence:

X<sup>1</sup>-Val-Cys-Ser-Glu-Gln-Ala-Glu-X<sup>2</sup>-Gly-Pro-  
Cys-Arg-Ala-X<sup>3</sup>-X<sup>4</sup>-X<sup>5</sup>-X<sup>6</sup>-Arg-Trp-Tyr-Phe-  
Asp-Val-Thr-Glu-Gly-Lys-Cys-Ala-Pro-Phe-  
Phe-Tyr-Gly-Gly-Cys-X<sup>7</sup>-Gly-Asn-Arg-Asn-  
Asn-Phe-Asp-Thr-Glu-Glu-Tyr-Cys-Met-Ala-  
Val-Cys-Gly-Ser-Ala-Ile,

wherein:

X<sup>1</sup> is selected from Glu-Val-Val-Arg-Glu-, Asp, or Glu;

X<sup>2</sup> is selected from Thr, Val, Ile and Ser;

X<sup>3</sup> is selected from Arg, Ala, Leu, Gly, or Met;

X<sup>4</sup> is selected from Ile, His, Leu, Lys, Ala, or Phe;

X<sup>5</sup> is selected from Ser, Ile, Pro, Phe, Tyr, Trp, Asn, Leu, His, Lys, or Glu;

X<sup>6</sup> is selected from Arg, His, or Ala; and

X<sup>7</sup> is selected from Gly, Ala, Lys, Pro, Arg, Leu, Met, or Tyr.

69. A protease inhibitor according to claim 68, wherein at least two amino acid residues selected from the group consisting of X<sup>3</sup>, X<sup>4</sup>, X<sup>5</sup>, and X<sup>6</sup> differ from the residues found in the naturally occurring sequence of KPI.

70. A protease inhibitor according to claim 68, wherein X<sup>1</sup> is Glu-Val-Val-Arg-Glu-, X<sup>2</sup> is Thr, Val, or Ser, X<sup>3</sup> is Ala or Leu, X<sup>4</sup> is Ile, X<sup>5</sup> is Tyr, X<sup>6</sup> is His and X<sup>7</sup> is Gly.

71. A protease inhibitor according to claim 70, wherein X<sup>2</sup> is Thr, and X<sup>3</sup> is Ala.

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72. A protease inhibitor according to claim 70, wherein X<sup>2</sup> is Thr, and X<sup>3</sup> is Leu.

73. A protease inhibitor according to claim 70, wherein X<sup>2</sup> is Val, and X<sup>3</sup> is Ala.

74. A protease inhibitor according to claim 70, wherein X<sup>2</sup> is Ser, and X<sup>3</sup> is Ala.

75. A protease inhibitor according to claim 70, wherein X<sup>2</sup> is Val, and X<sup>3</sup> is Leu.

76. A protease inhibitor according to claim 70, wherein X<sup>2</sup> is Ser, and X<sup>3</sup> is Leu.

77. A protease inhibitor according to claim 68, wherein X<sup>1</sup> is Glu-Val-Val-Arg-Glu-, X<sup>2</sup> is Thr, X<sup>3</sup> is Leu, X<sup>4</sup> is Phe, X<sup>5</sup> is Lys, X<sup>6</sup> is Arg and X<sup>7</sup> is Gly.

78. A protease inhibitor according to claim 68, wherein X<sup>1</sup> is Glu-Val-Val-Arg-Glu-, X<sup>2</sup> is Thr, X<sup>3</sup> is Leu, X<sup>4</sup> is Phe, X<sup>5</sup> is Lys, X<sup>6</sup> is Arg and X<sup>7</sup> is Tyr.

79. A protease inhibitor according to claim 68, wherein X<sup>1</sup> is Glu-Val-Val-Arg-Glu-, X<sup>2</sup> is Thr, X<sup>3</sup> is Leu, X<sup>4</sup> is Phe, X<sup>5</sup> is Lys, X<sup>6</sup> is Arg and X<sup>7</sup> is Leu.